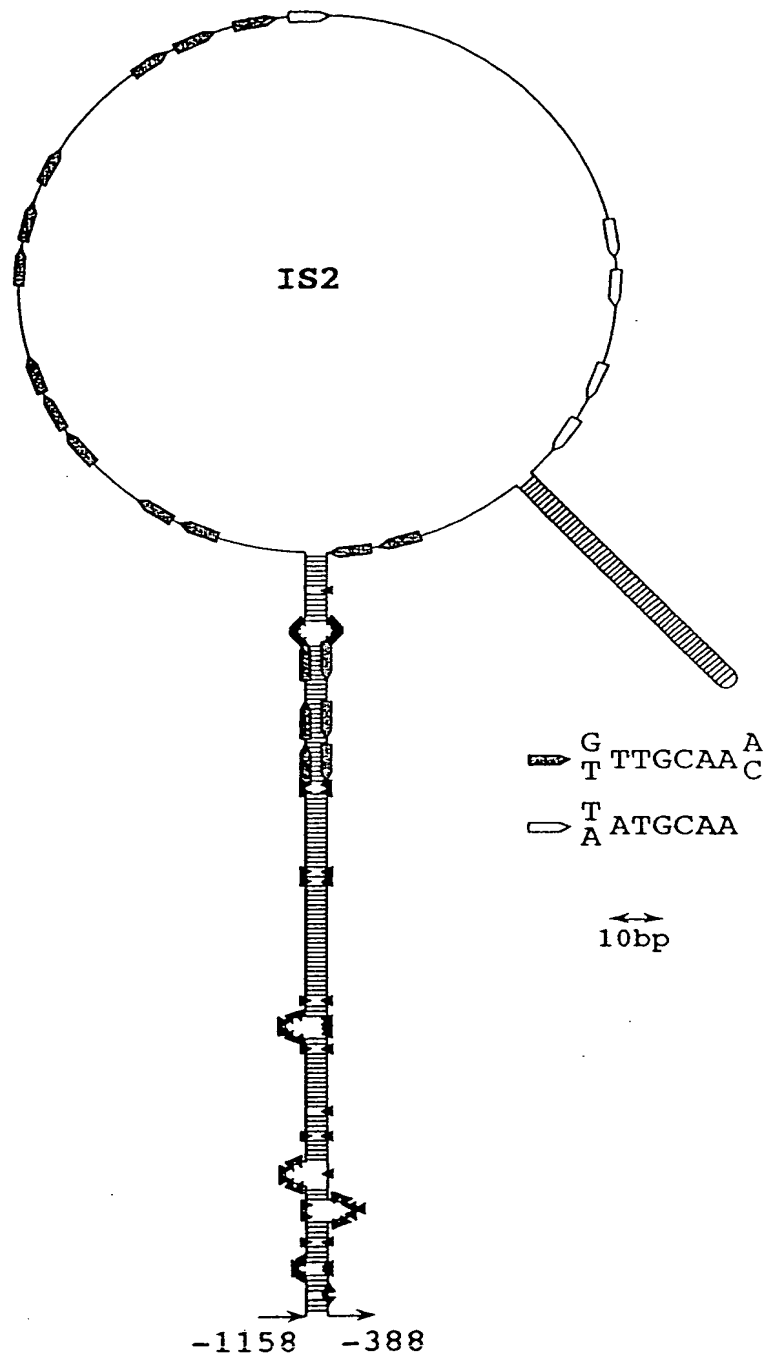


FIG. 1



[illegible]

FIG. 3

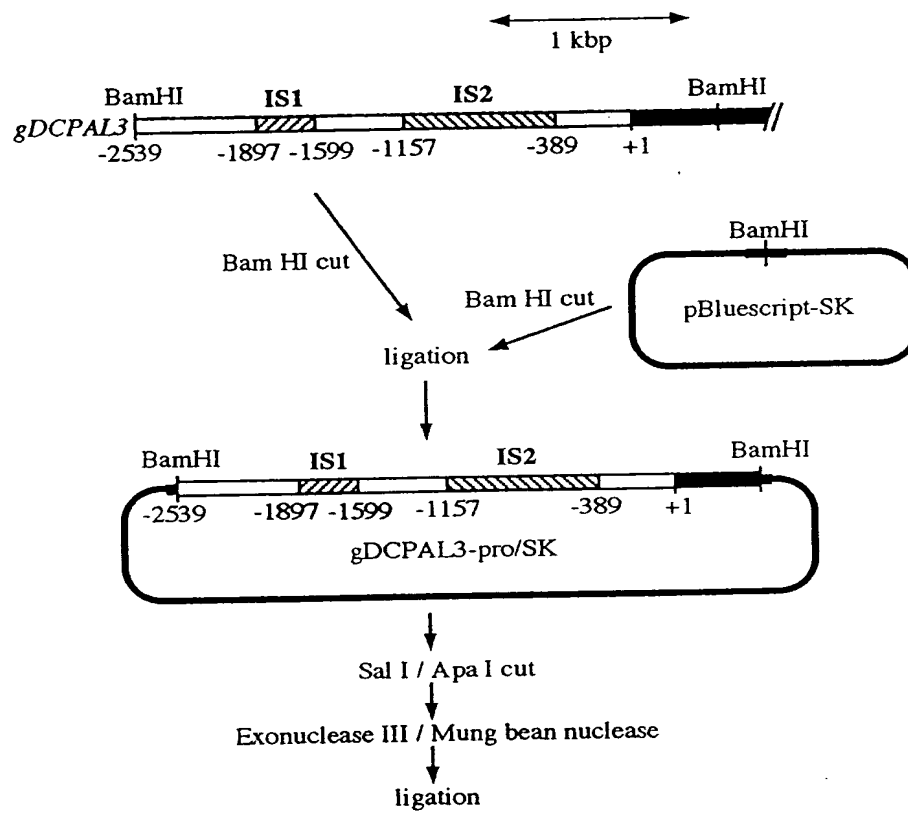


FIG. 4

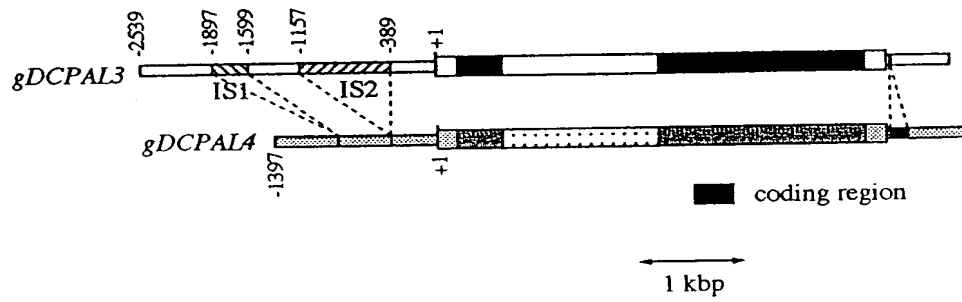


FIG. 5

Stowaway

Gene	Position		size(bp)
IS1 \square	(5'-1597)	TACTCCCTACGTCGCCATTTTACGACCGTCATTTTCTTTT	303
carrot DC8	(5'-514)	TACTCTCTCTATCTCATTTTAACTCTATACA. TTTCTTTT	253
carrot DC59	(5'-409)	TACTCCCTTCCCTACCTTTTCACTTATCAGTTTCTATTT	249
petunia CH8	(in1 884)	TACTCCCTTCCCTCCCTTTTATTTGATG. TTTTCTCTTT	122
tomato PAL	(5'-494)	TACTCTCTCTCTTCTTAAAGAAATGCTTATTTCTTTT	248
parsley PR2	(5'-415)	TACTCCCTTCCCTTCTTAAAGAAATGCTTATTTCTTTT	243
tobacco CHM	(5'-209)	TACTCCCTTCCCTTCTTAAAGAAATGCTTATTTCTTTT	244
pea lectin	(5'-1296)	AACTCCCTTCCCTTCTTAAAGAAATGCTTATTTCTTTT	275
rape EP8P	(in5 1835)	TACTCCCTTCTTCTTAAAGAAATGCTTATTTCTTTT	220
potato patatin	(in2 922)	TACTCTCTTCTTCTTAAAGAAATGCTTATTTCTTTT	259
rice PCMA	(5'-432)	TACTCTCTCTCTTCTTAAAGAAATGCTTATTTCTTTT	122
maize xein	(3' UTR)	TACTCCCTTCTCTTCTTAAAGAAATGCTTATTTCTTTT	163
wheat AMY	(5'-458)	TACTCTCTCTCTTCTTAAAGAAATGCTTATTTCTTTT	100
barley RcaA	(in3 839)	TACTCCCTTCTCTTCTTAAAGAAATGCTTATTTCTTTT	159
sugarcane PEPC	(in6 2675)	TACTCCCTTCTCTTCTTAAAGAAATGCTTATTTCTTTT	267
sorghum PEPC	(5'-469)	TACTCCCTTCTCTTCTTAAAGAAATGCTTATTTCTTTT	255

Legend:

- m^7G (cap)
- A (adenine)
- U (uracil)
- G (guanine)
- C (cytosine)

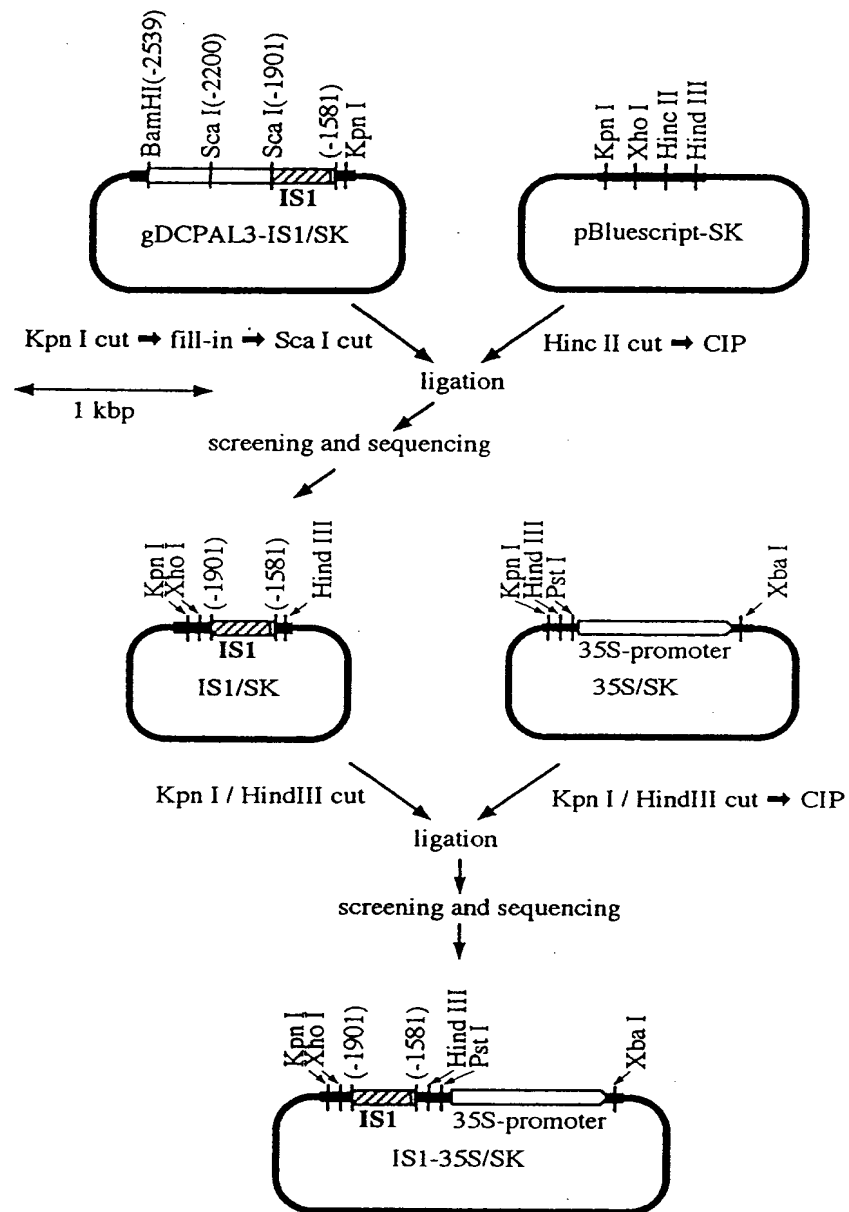
Scale bar: 10bp

Sequences and positions:

gDCPAL3 (black):
 5' ...
 -1174 ...
 -1158 ...
 -388 ...
 -360 ...

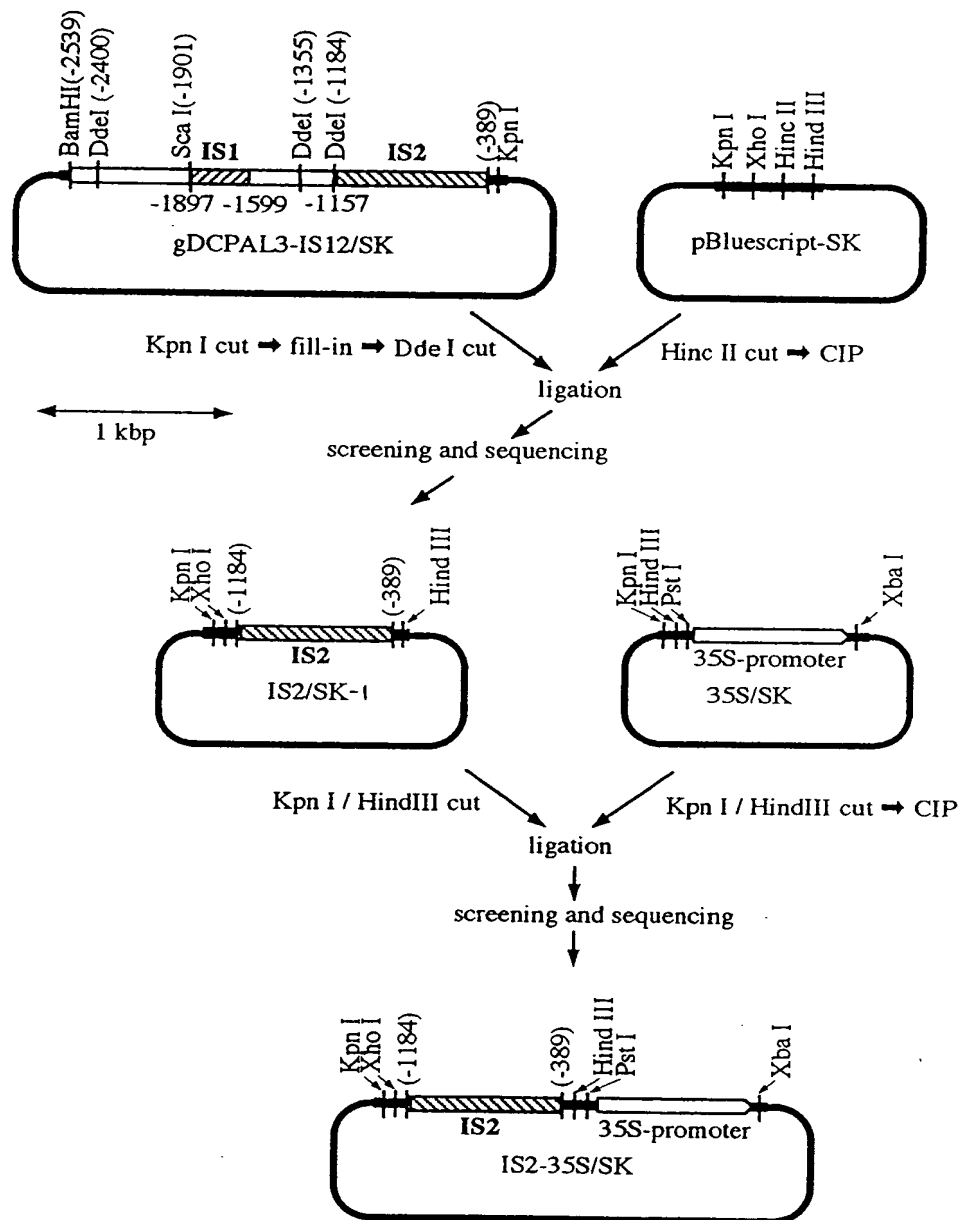
gDCPAL4 (grey):
 5' ...
 -396 ...
 -380 ...
 -379 ...
 -360 ...

FIG. 7



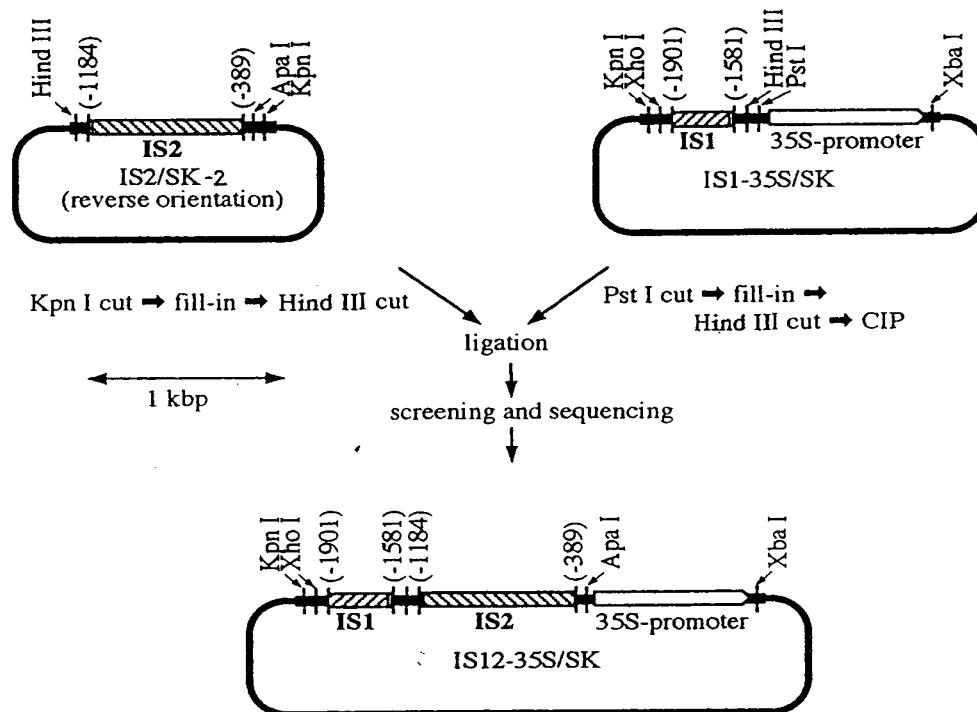
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FIG. 8



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FIG. 9



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FIG. 10

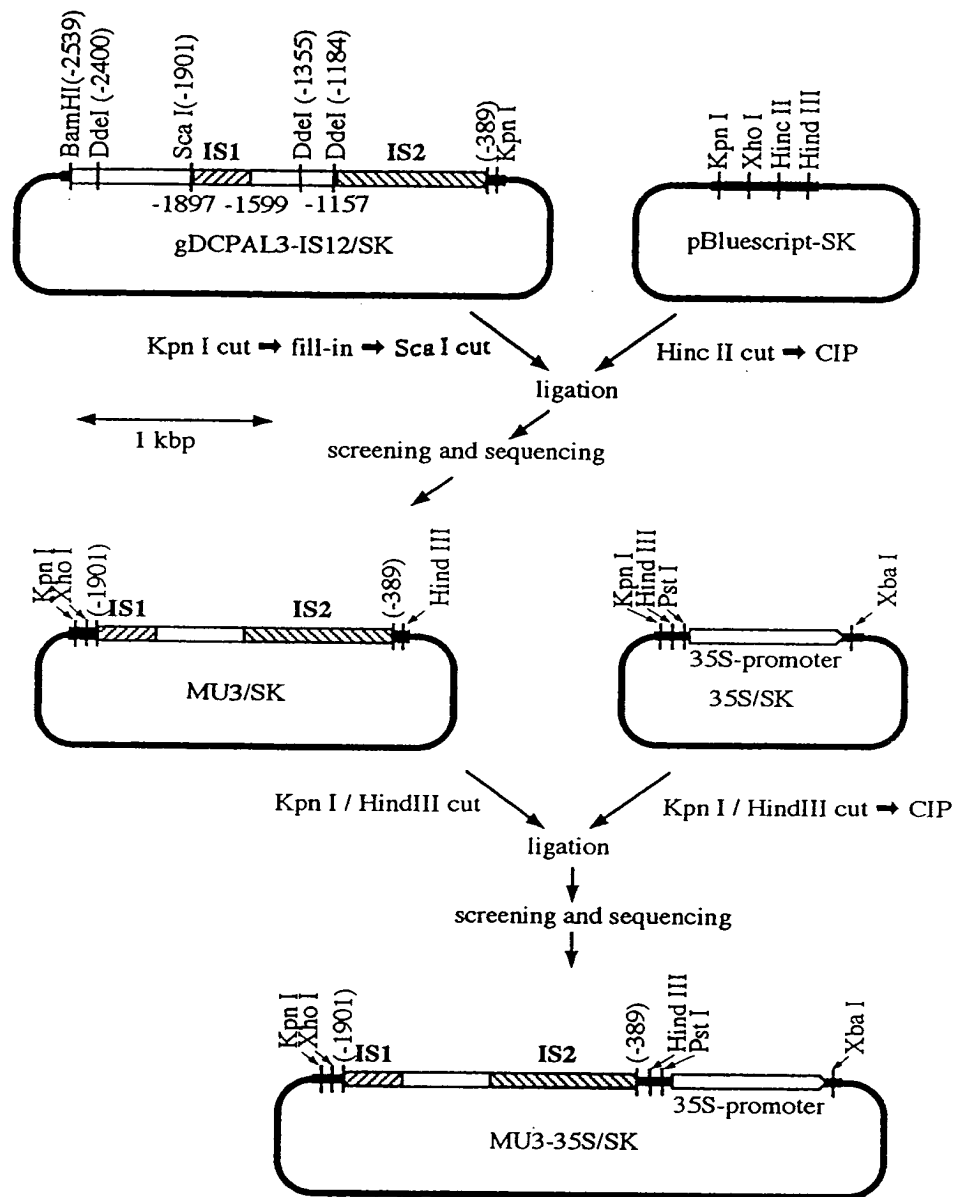
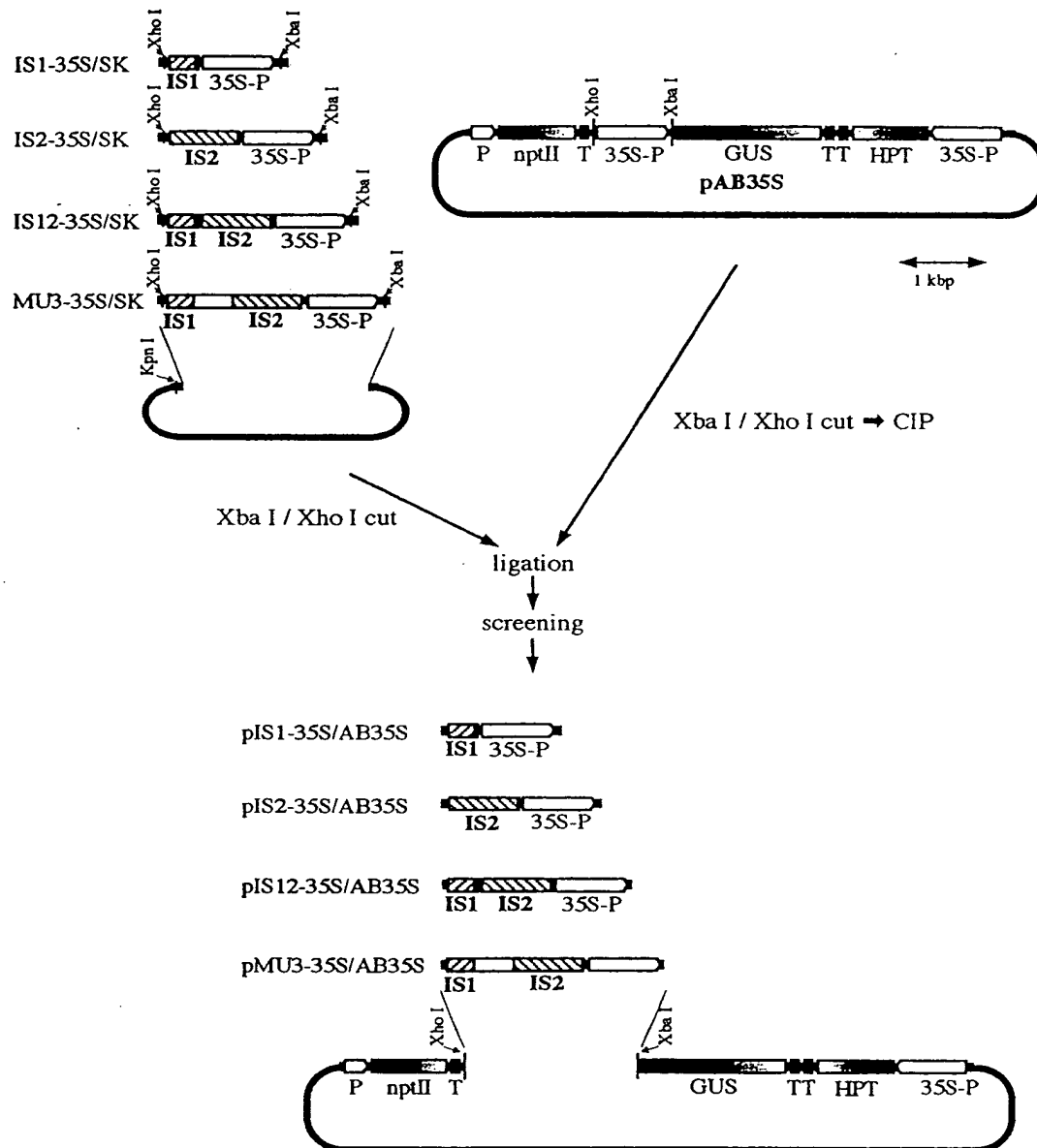
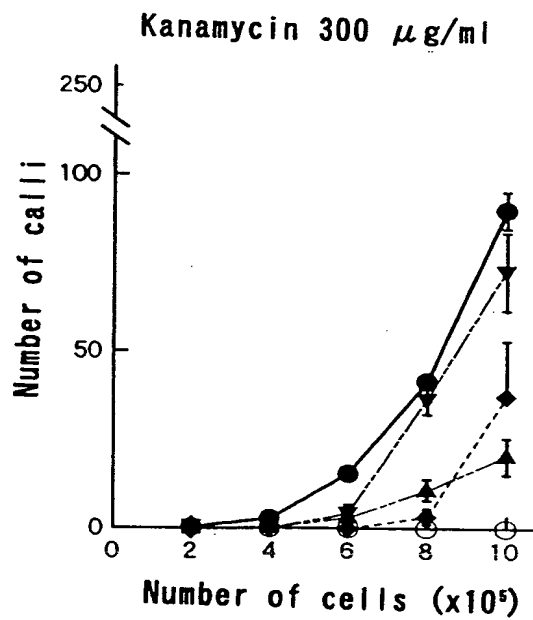
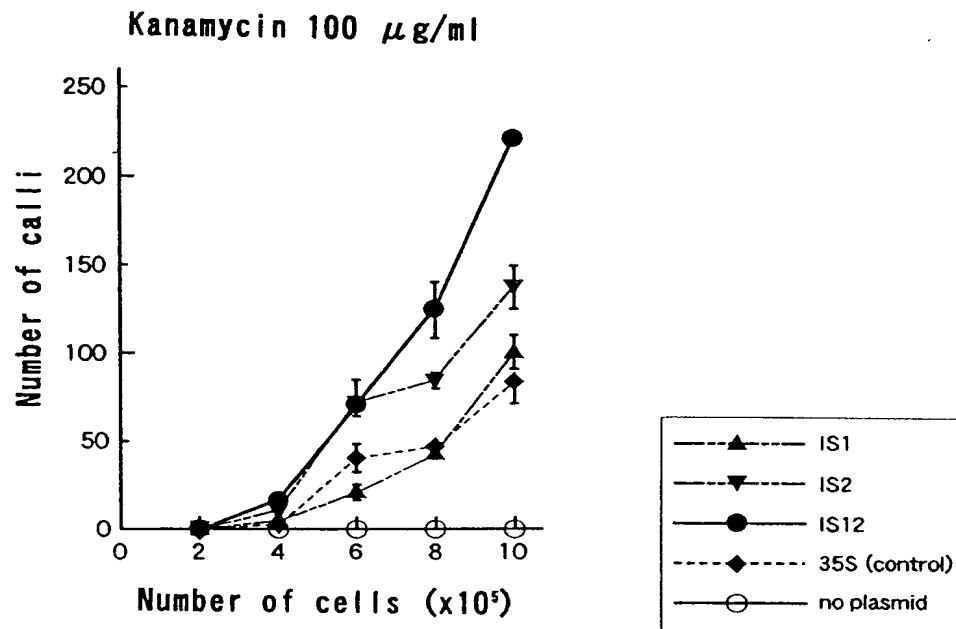


FIG. 11



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FIG. 12



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FIG. 13

